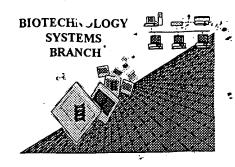
## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/9/0,033	
Source:	OIPE	
Date Processed by STIC:	7/30/201	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## **Checker Version 3.0**

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING DATE: 07/30/2001 PATENT APPLICATION: US/09/910,033 TIME: 16:03:42

Input Set : A:\210212US.txt

Output Set: N:\CRF3\07302001\I910033.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: RIEBEL, Bettina HUMMEL, Werner BOMMARIUS, Andreas 7 <120> TITLE OF INVENTION: RECOMBINANT ENZYMES HAVING IMPROVED NAD(H) ACCEPTANCE 9 <130> FILE REFERENCE: 210212US 11 <140> CURRENT APPLICATION NUMBER: US/09/910,033 11 <141> CURRENT FILING DATE: 2001-07-23 11 <150> PRIOR APPLICATION NUMBER: DE 10037101.9 12 <151> PRIOR FILING DATE: 2000-07-27 14 <160> NUMBER OF SEQ ID NOS: 7 16 <170> SOFTWARE: PatentIn version 3.1 18 <210> SEQ ID NO: 1 19 <211> LENGTH: 759 20 <212> TYPE: DNA 21 <213> ORGANISM: Lactobacillus brevis 23 <220> FEATURE: 24 <221> NAME/KEY: CDS 25 <222> LOCATION: (1)..(759) 26 <223> OTHER INFORMATION: 29 <400> SEQUENCE: 1 30 atg tct aac cgt ttg gat ggt aag gta gca atc att aca ggt ggt acg 48 31 Met Ser Asn Arg Leu Asp Gly Lys Val Ala Ile Ile Thr Gly Gly Thr 34 ttg qgt atc ggt tta gct atc gcc acg aag ttc gtt gaa gaa ggg gct 96 35 Leu Gly Ile Gly Leu Ala Ile Ala Thr Lys Phe Val Glu Glu Gly Ala 20 25 144 38 aag gtc atg att acc gac cgg cac agc gat gtt ggt gaa aaa gca gct 39 Lys Val Met Ile Thr Asp Arg His Ser Asp Val Gly Glu Lys Ala Ala 35 42 aag agt gtc ggc act cct gat cag att caa ttt ttc caa cat gat tct 192 43 Lys Ser Val Gly Thr Pro Asp Gln Ile Gln Phe Phe Gln His Asp Ser 46 too gat gaa gao ggo tgg aog aaa tta tto gat goa aog gaa aaa goo 240 47 Ser Asp Glu Asp Gly Trp Thr Lys Leu Phe Asp Ala Thr Glu Lys Ala 70 75 50 ttt ggc cca gtt tct aca tta gtt aat aac gct ggg atc gcg gtt aac 288 51 Phe Gly Pro Val Ser Thr Leu Val Asn Asn Ala Gly Ile Ala Val Asn 54 aag agt gtc gaa gaa acc acg act gct gaa tgg cgt aaa tta tta gcc 336 55 Lys Ser Val Glu Glu Thr Thr Ala Glu Trp Arg Lys Leu Leu Ala 100 105 384 58 gtc aac ctt gat ggt gtc ttc ttc ggt acc cga tta ggg att caa cgg 59 Val Asn Leu Asp Gly Val Phe Phe Gly Thr Arg Leu Gly Ile Gln Arg 115 120 62 atg aag aac aaa ggc tta ggg gct tcc atc atc aac atg tct tcg atc 432 63 Met Lys Asn Lys Gly Leu Gly Ala Ser Ile Ile Asn Met Ser Ser Ile 64 130 135 140

RAW SEQUENCE LISTING DATE: 07/30/2001 PATENT APPLICATION: US/09/910,033 TIME: 16:03:42

Input Set : A:\210212US.txt

Output Set: N:\CRF3\07302001\I910033.raw

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68						150					155					160	
		_	-			atg		_		_	_		_				528
71 72	GIY	Ala	Val	Arg	11e 165	Met	Ser	Lys	Ser	A1a 170	Ala	Leu	Asp	Cys	A1a 175	Leu	
						cgg											576
	Lys	Asp	Tyr	Asp	Val	Arg	Val	Asn		Val	His	Pro	Gly		Ile	Lys	
76				180					185					190			
			_	-	_	gac				_	-	-		_			624
79 80	Thr	Pro	195	vaı	ASP	Asp	Leu	200	СТА	Ата	GIU	Glu	205	met	ser	GIN	
82	cgg	acc	aag	acg	cca	atg	ggc	cat	atc	ggt	gaa	cct	aac	gat	att	gcc	672
						Met											
84		210					215					220					
			-	-		ttg	-			-							720
	_	Ile	Cys	Val	Tyr	Leu	Ala	Ser	Asn	Glu		Lys	Phe	Ala	Thr		
88						230					235					240	750
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91 92	ser	GLU	Pne	vaı	vai 245	Asp	GIY	GIY	Tyr	250	Ата	GIN					
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102	Met	Ser	Asn	Arg	Leu	. Asp	Gly	Lys	Val	. Ala	ı Ile	: Ile	Thr	Gly	Gly	Thr	
103					5					10					15		
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115	_	50	, 41	. 017		110	55	0111		. 011		60	. 011		, ,,,,,,		
			Glu	Asp	Gly	Trp		Lys	Leu	Phe	Asp		Thr	Glu	Lys	Ala	
119		-		-	-	70		-			75				_	80	
122	Phe	Gly	Pro	Val	Ser	Thr	Leu	Val	Asn	Asn	Ala	Gly	, Ile	Ala	Val	Asn	
123																	
					85					90					95		
126	Lys	Ser	Val	Glu		Thr	Thr	Thr	Ala					Leu		Ala	
127				100	Glu				105	Glu	Trp	Arg	Lys	110	Leu		
127 130	Val		Leu	100 Asp	Glu			Phe	105 Gly	Glu	Trp	Arg	Lys Gly	110	Leu	Ala	
127 130 131	Val	Asn	Leu 115	100 Asp	Glu Gly	Val	Phe	Phe	105 Gly	Glu	Trp	Arg	Lys Gly 125	110 Ile	Leu Gln	Arg	
127 130 131 134	Val Met	Asn Lys	Leu 115 Asn	100 Asp	Glu Gly	Val	Phe Gly	Phe 120 Ala	105 Gly	Glu	Trp	Arg Leu Asn	Lys Gly 125 Met	110 Ile	Leu Gln		
127 130 131 134 135	Val Met	Asn Lys 130	Leu 115 Asn	100 Asp Lys	Glu Gly Gly	Val	Phe Gly 135	Phe 120 Ala	105 Gly Ser	Glu Thr	Trp	Arg Leu Asn 140	Lys Gly 125 Met	110 Ile	Leu Gln	Arg	
127 130 131 134 135 138	Val Met Glu	Asn Lys 130 Gly	Leu 115 Asn	100 Asp Lys	Glu Gly Gly	Val	Phe Gly 135	Phe 120 Ala	105 Gly Ser	Glu Thr	Trp	Leu Asn 140	Lys Gly 125 Met	110 Ile	Leu Gln	Arg	
127 130 131 134 135 138 139	Val Met Glu 145	Asn Lys 130 Gly	Leu 115 Asn Phe	100 Asp Lys	Glu Gly Gly Gly	Val Leu Asp 150	Phe Gly 135 Pro	Phe 120 Ala Ser	105 Gly Ser	Glu Thr Ile	Trp Arg	Leu Asn 140	Lys Gly 125 Met	110 Ile Ser	Leu Gln Ser	Arg Lys 160	
127 130 131 134 135 138 139	Val Met Glu 145 Gly	Asn Lys 130 Gly	Leu 115 Asn Phe	100 Asp Lys	Glu Gly Gly Gly	Val Leu Asp 150 Met	Phe Gly 135 Pro	Phe 120 Ala Ser	105 Gly Ser	Glu Thr Ile	Trp Arg	Leu Asn 140	Lys Gly 125 Met	110 Ile Ser	Leu Gln Ser	Arg Lys 160 Leu	
127 130 131 134 135 138 139 142	Val Met Glu 145 Gly	Asn Lys 130 Gly Ala	Leu 115 Asn Phe	100 Asp Lys Val	Glu Gly Gly Gly Ile	Val Leu Asp 150 Met	Phe Gly 135 Pro	Phe 120 Ala Ser	105 Gly Ser Leu Ser	Thr Ile Gly Ala	Trp Arg	Leu Asn 140 Tyr	Glys 125 Met	110 Ile Ser Ala	Leu Glm Ser Ser Ala 175	Arg Lys 160 Leu	

DATE: 07/30/2001

PATENT APPLICATION: US/09/910,033 TIME: 16:03:42 Input Set : A:\210212US.txt Output Set: N:\CRF3\07302001\I910033.raw 185 190 180 150 Thr Pro Leu Val Asp Asp Leu Pro Gly Ala Glu Glu Ala Met Ser Gln 200 195 154 Arg Thr Lys Thr Pro Met Gly His Ile Gly Glu Pro Asn Asp Ile Ala 215 158 Tyr Ile Cys Val Tyr Leu Ala Ser Asn Glu Ser Lys Phe Ala Thr Gly 230 235 162 Ser Glu Phe Val Val Asp Gly Gly Tyr Thr Ala Gln 163 245 166 <210> SEQ ID NO: 3 167 <211> LENGTH: 24 168 <212> TYPE: DNA 169 <213> ORGANISM: Artificial Sequence 171 <220> FEATURE: 172 <223> OTHER INFORMATION: synthetic DNA 174 <400> SEQUENCE: 3 24 175 accgaccggc acagcgatgt tggt 178 <210> SEQ ID NO: 4 179 <211> LENGTH: 8 180 <212> TYPE: PRT 181 <213> ORGANISM: Artificial Sequence 183 <220> FEATURE: 184 <223> OTHER INFORMATION: synthetic peptide 186 <400> SEQUENCE: 4 188 Thr Asp Arg His Ser Asp Val Gly 189 1 192 <210> SEQ ID NO: 5 193 <211> LENGTH: 24 194 <212> TYPE: DNA 195 <213> ORGANISM: Artificial Sequence 197 <220> FEATURE: 198 <223> OTHER INFORMATION: synthetic DNA 200 <400> SEQUENCE: 5 201 accaacatcg ctgtgccggt cggt 24 204 <210> SEQ ID NO: 6 205 <211> LENGTH: 8 206 <212> TYPE: DNA This is not a pertile sequence. 207 <213> ORGANISM: Artificial Sequence 209 <220> FEATURE: 210 <223> OTHER INFORMATION: synthetic peptide 212 <400> SEQUENCE: 6 213 gvdshrdt 216 <210> SEQ ID NO: 7 217 <211> LENGTH: 8 218 <212> TYPE: DNA 219 <213> ORGANISM: Artificial Sequence 222 <223 OTHER INFORMATION: synthetic (peptide) hot a peptide requese 221 <220> FEATURE:

RAW SEQUENCE LISTING

224 <400> SEQUENCE: 7

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/910,033

DATE: 07/30/2001

TIME: 16:03:42

Input Set : A:\210212US.txt
Output Set: N:\CRF3\07302001\I910033.raw

225 tdrhsdvg

8

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/910,033

DATE: 07/30/2001 TIME: 16:03:43

Input Set : A:\210212US.txt
Output Set: N:\CRF3\07302001\I910033.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date